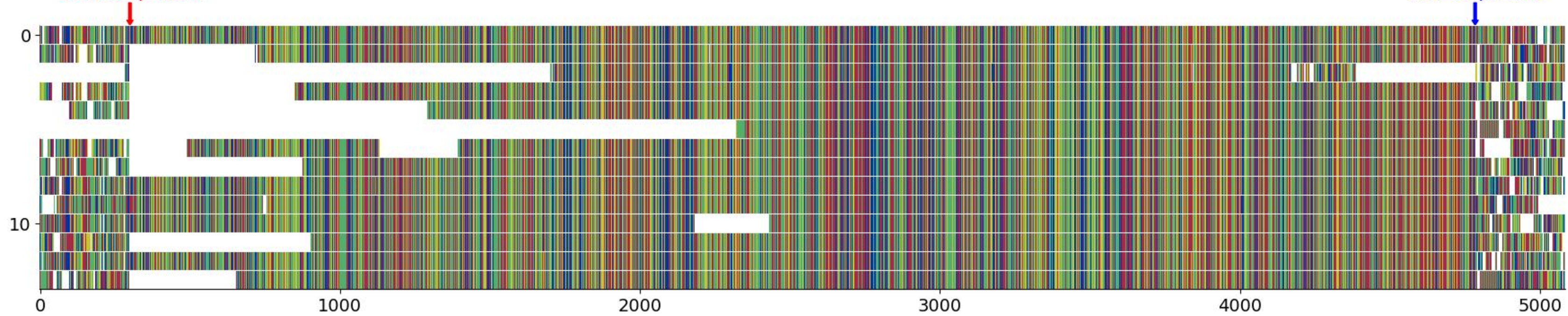


MSA length = 4485

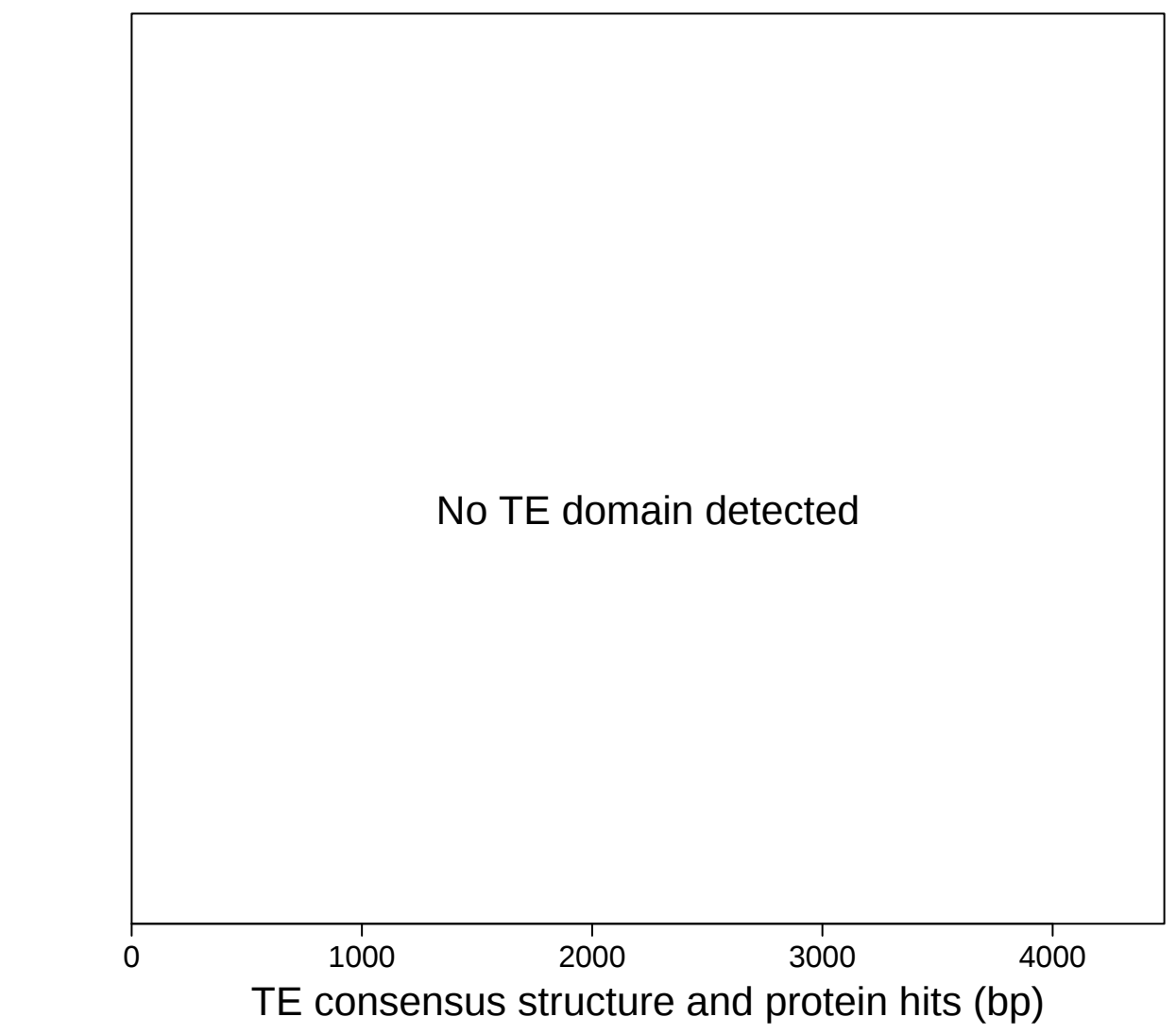
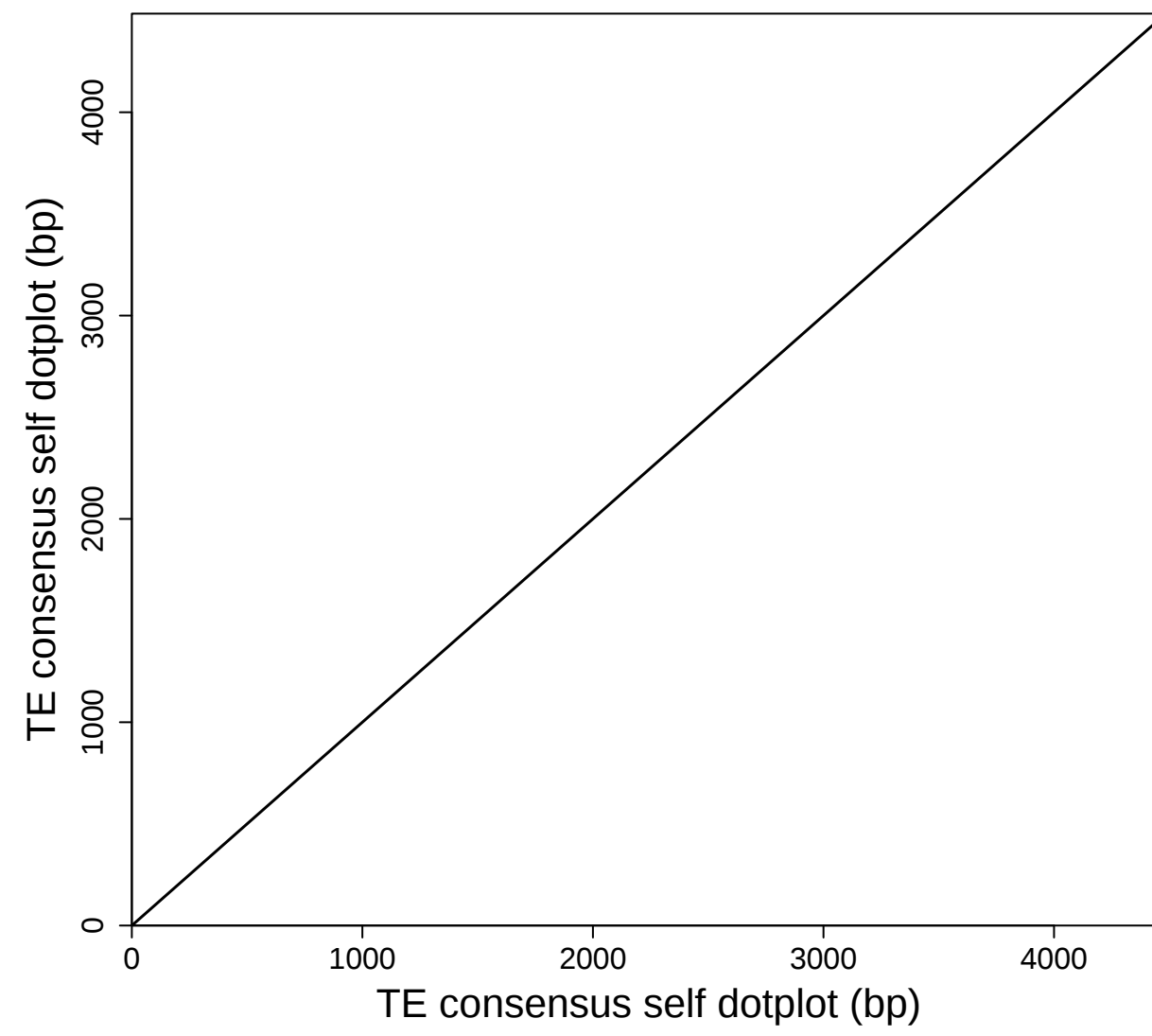
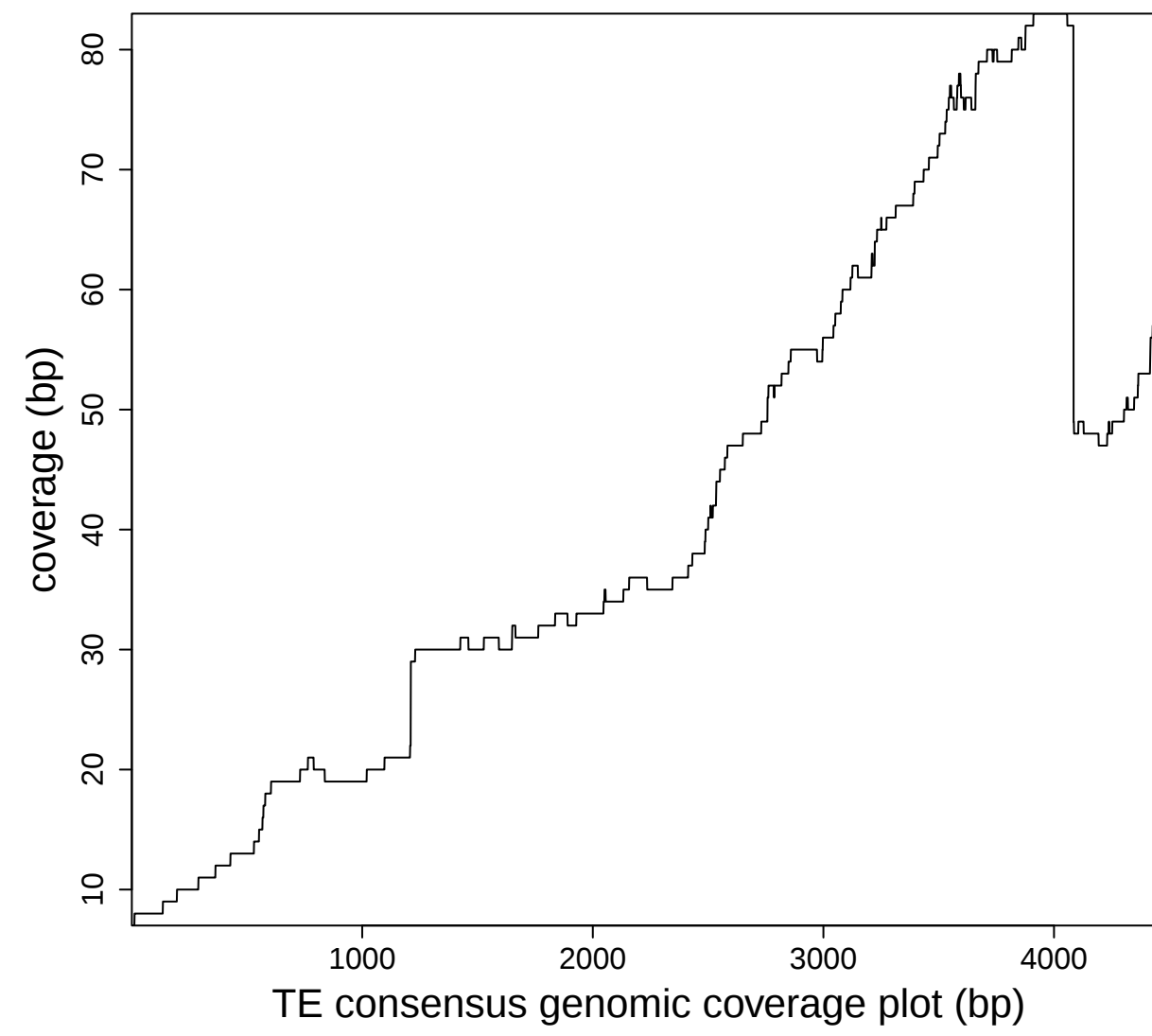
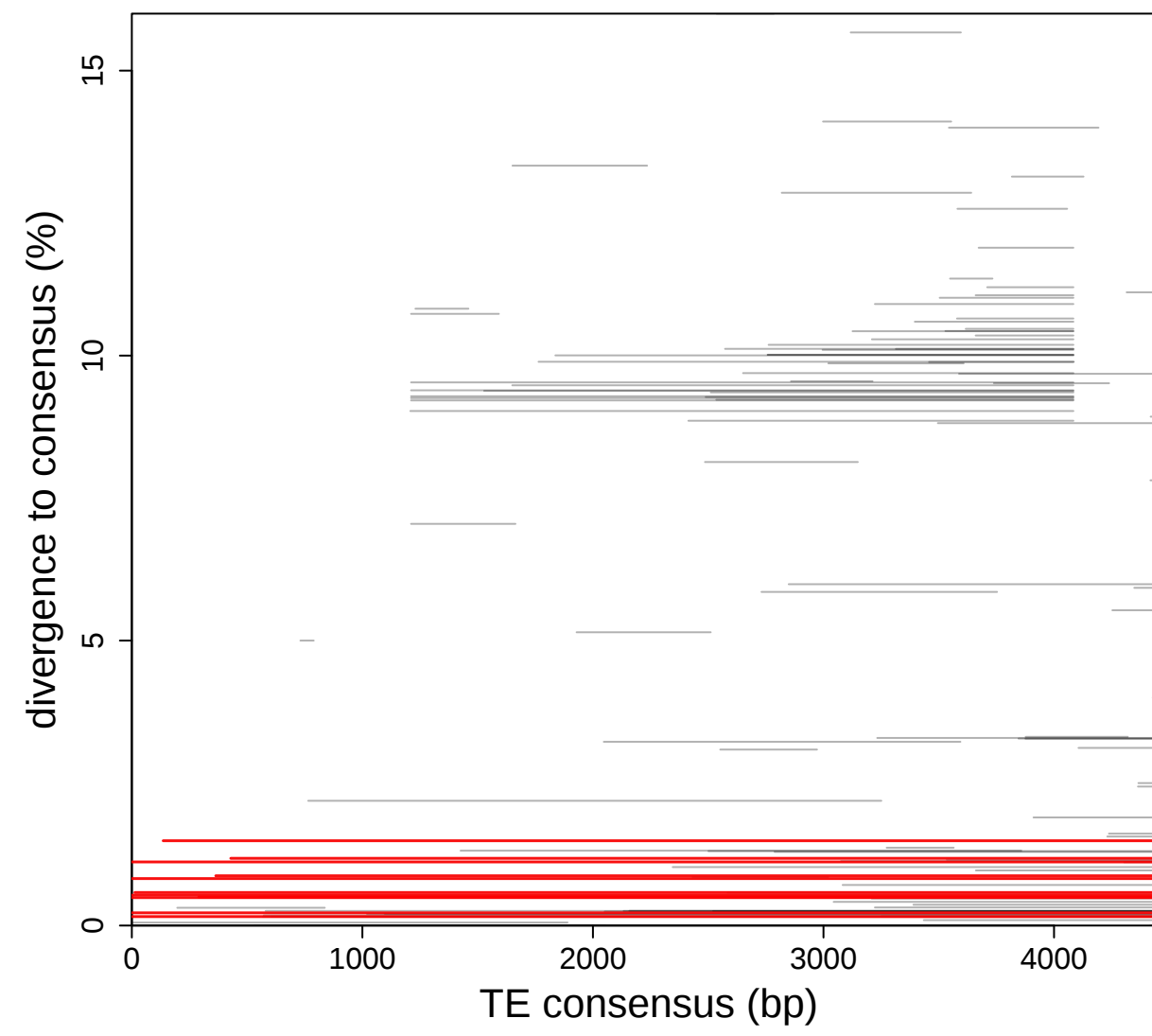
Start crop Point

End crop Point



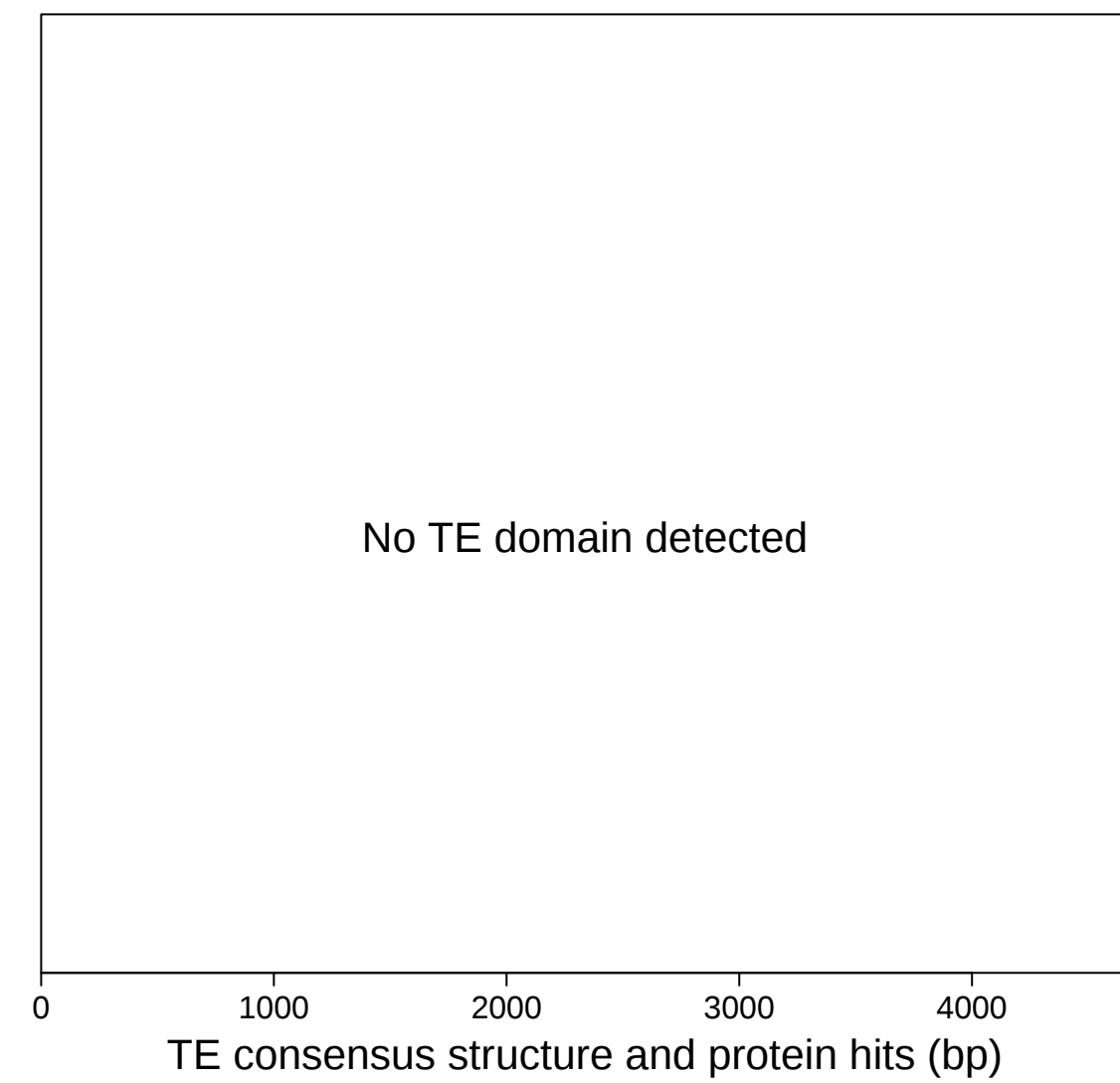
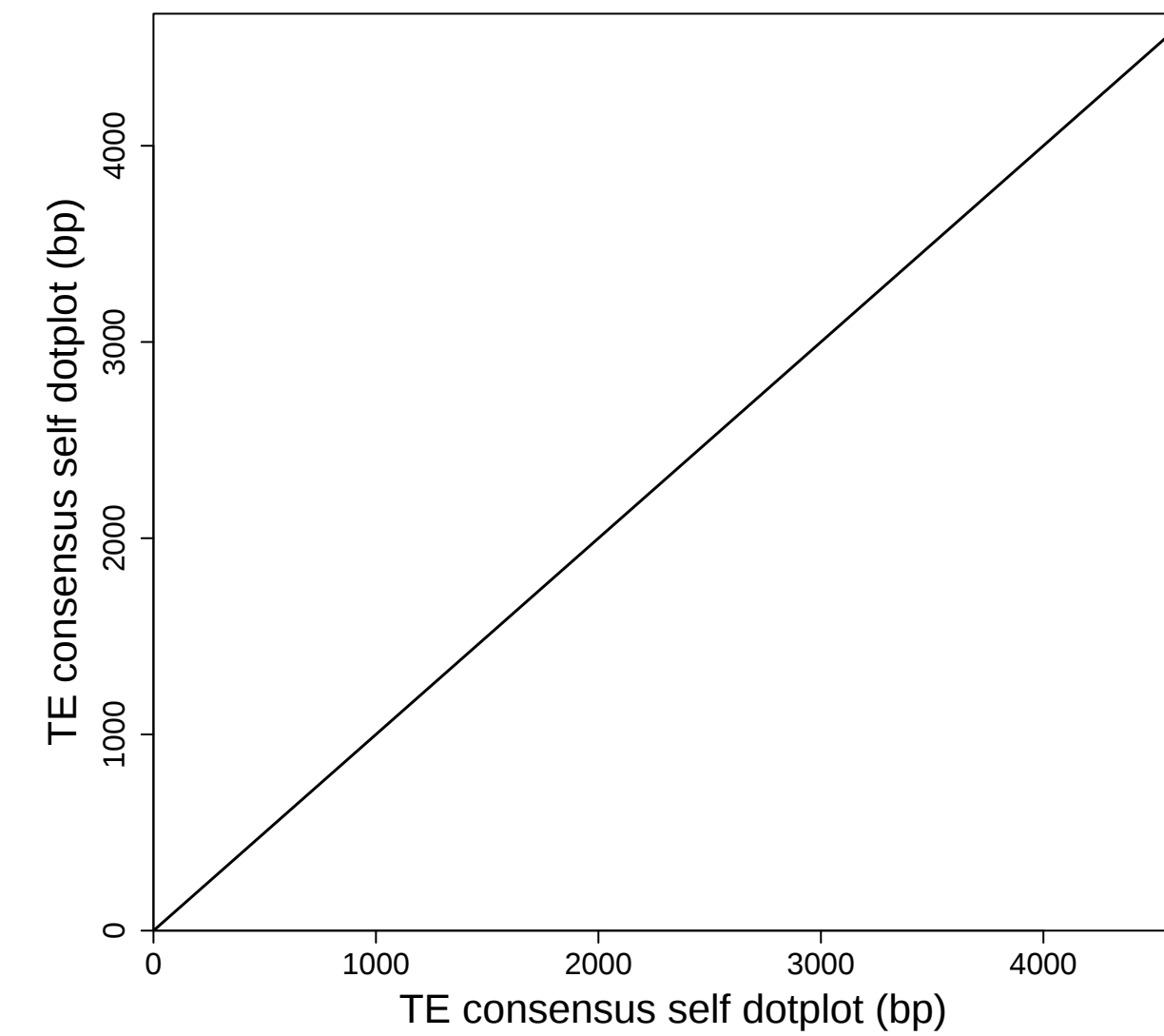
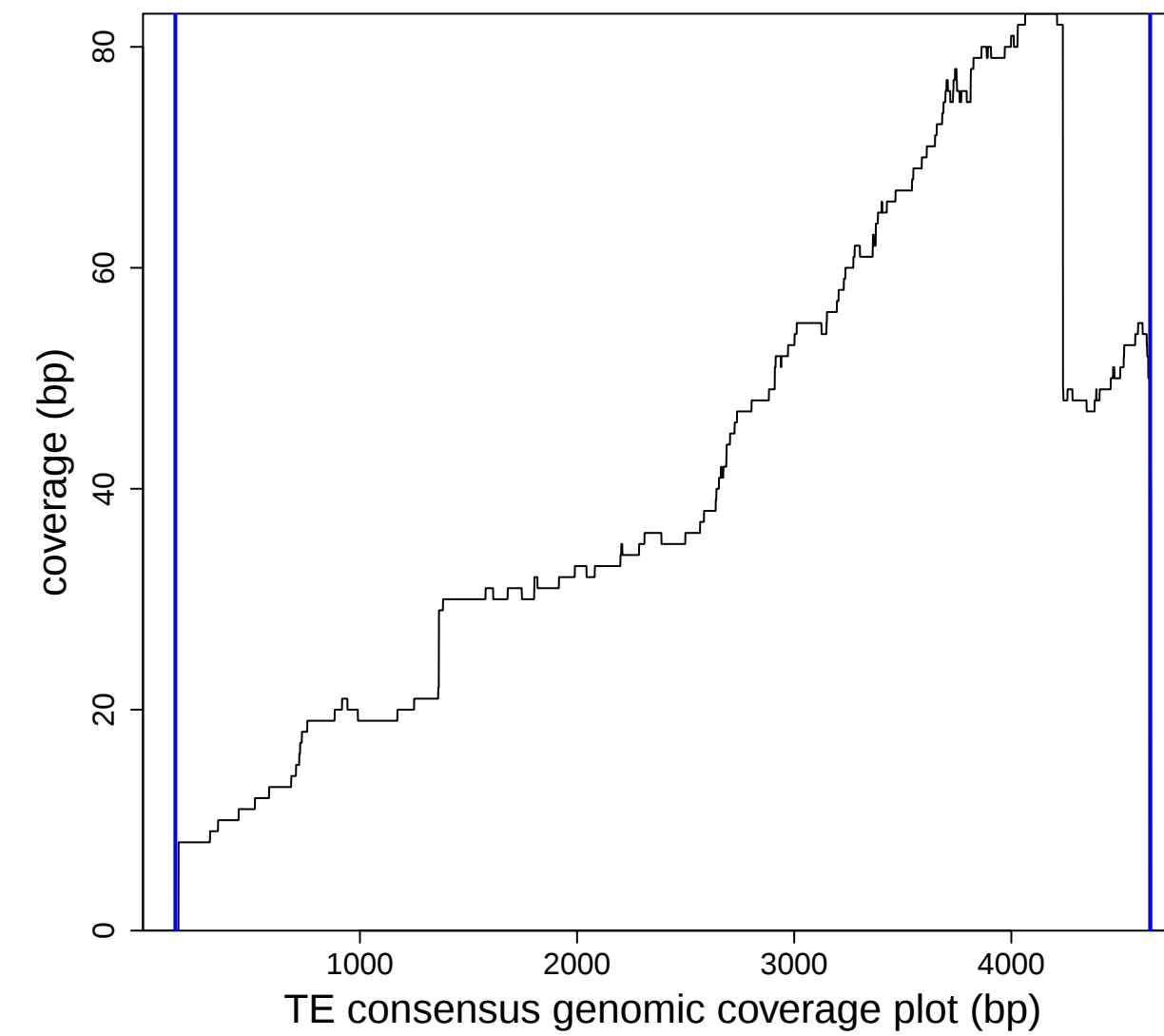
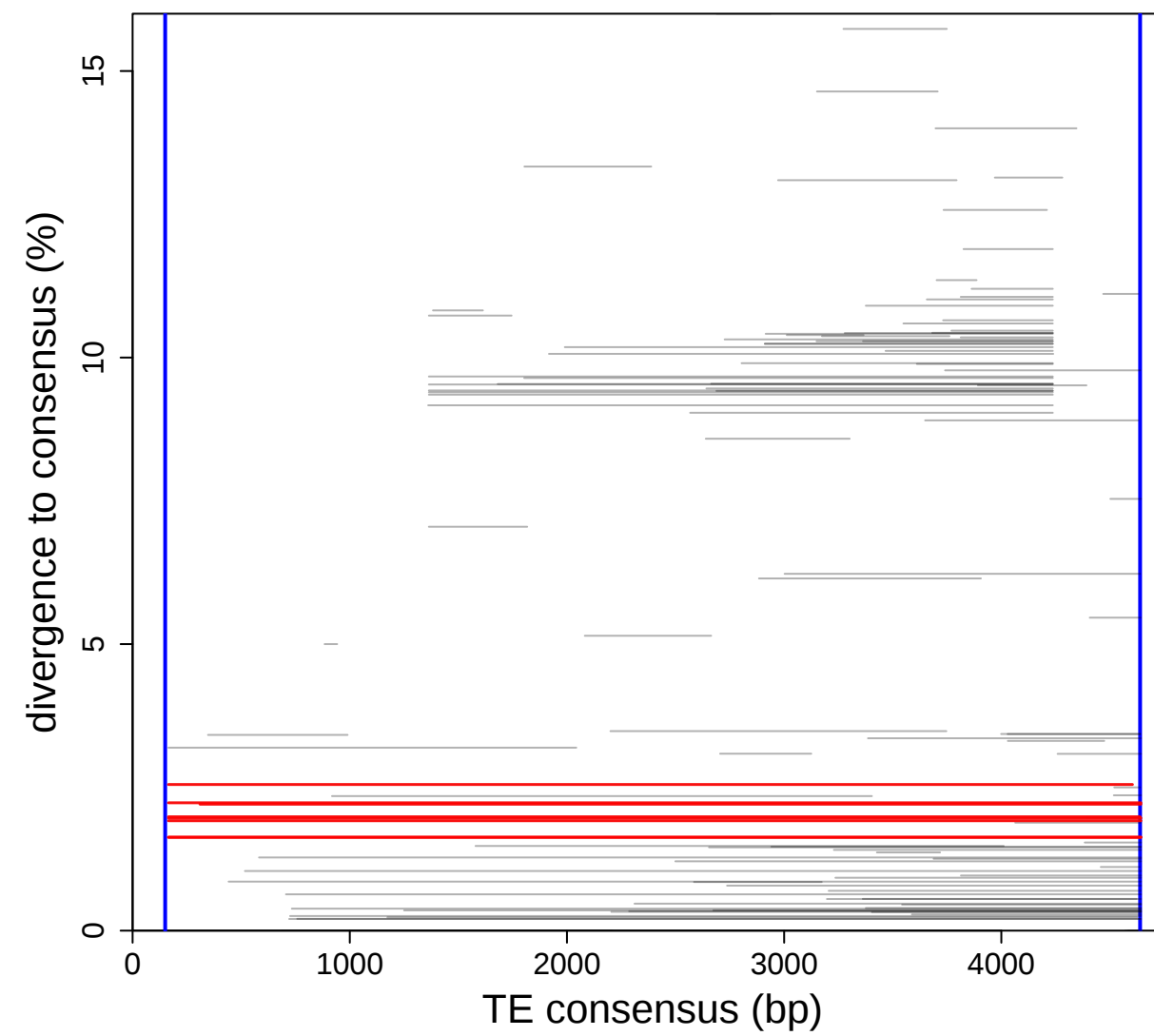
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 4485bp; fragments: 124; full length: 11 (≥ 4036.5 bp)

After TEtrimmer 4485 bp

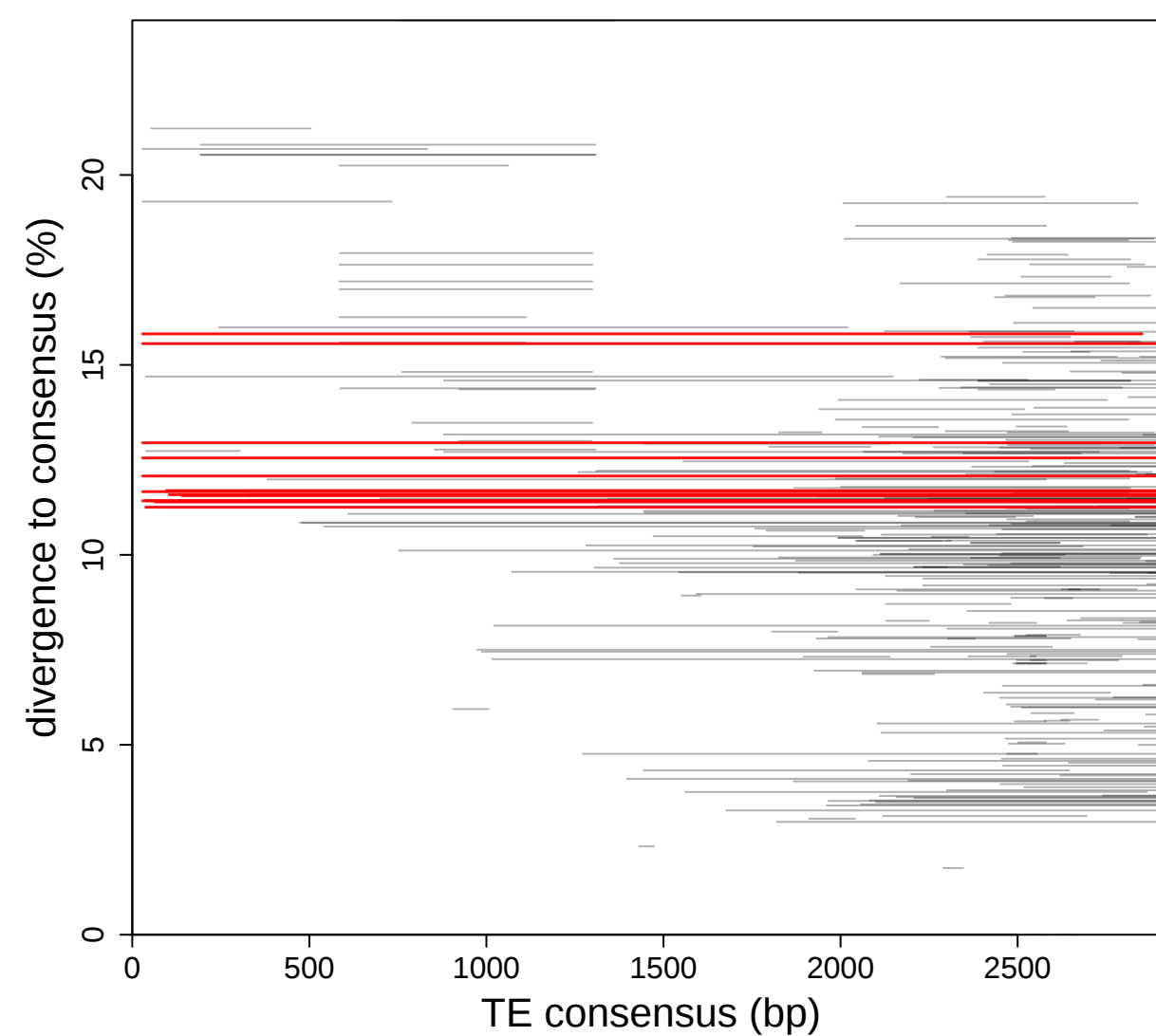


3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.fa
size: 4789bp; fragments: 118; full length: 8 (>=4310.1bp)

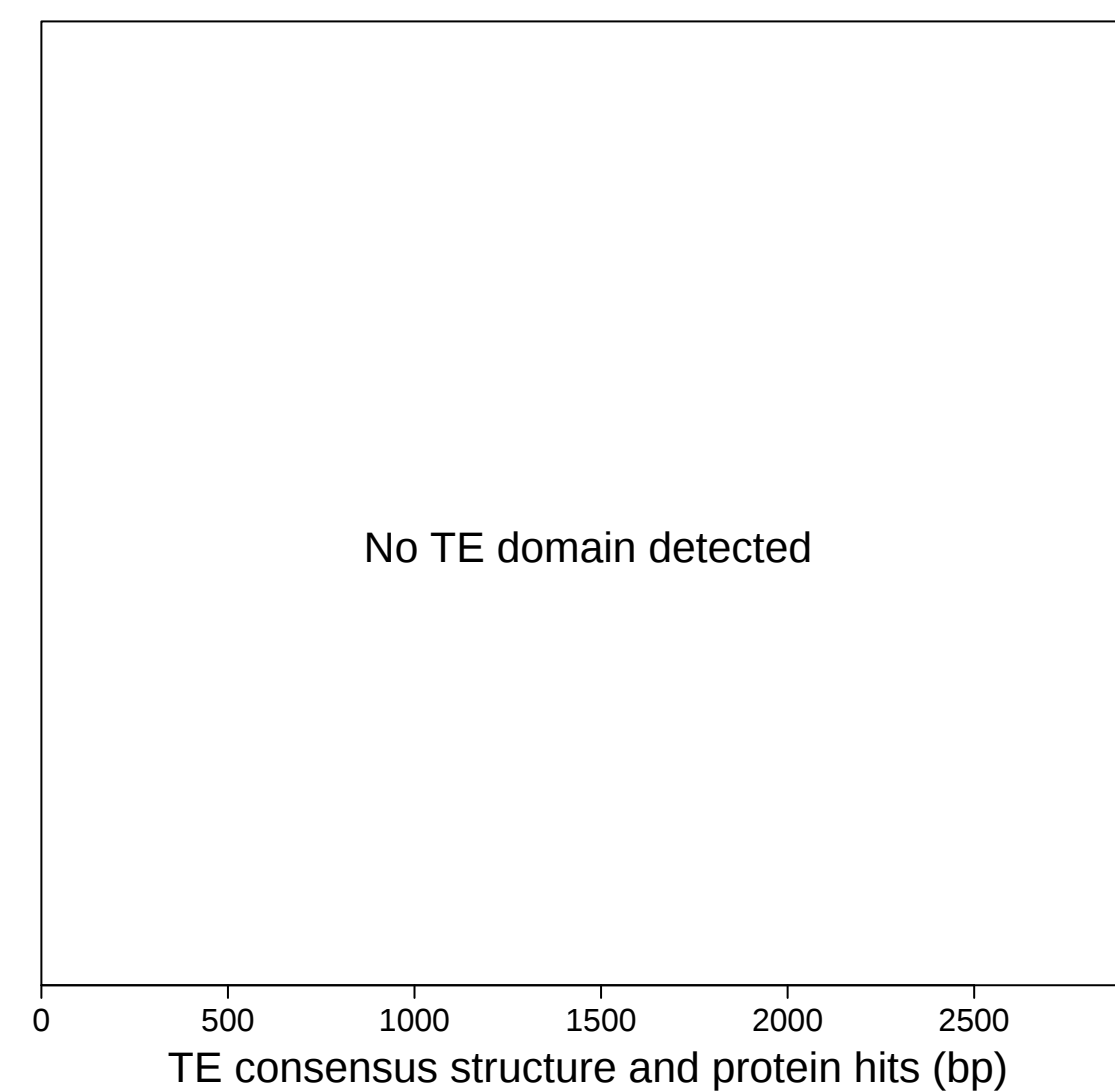
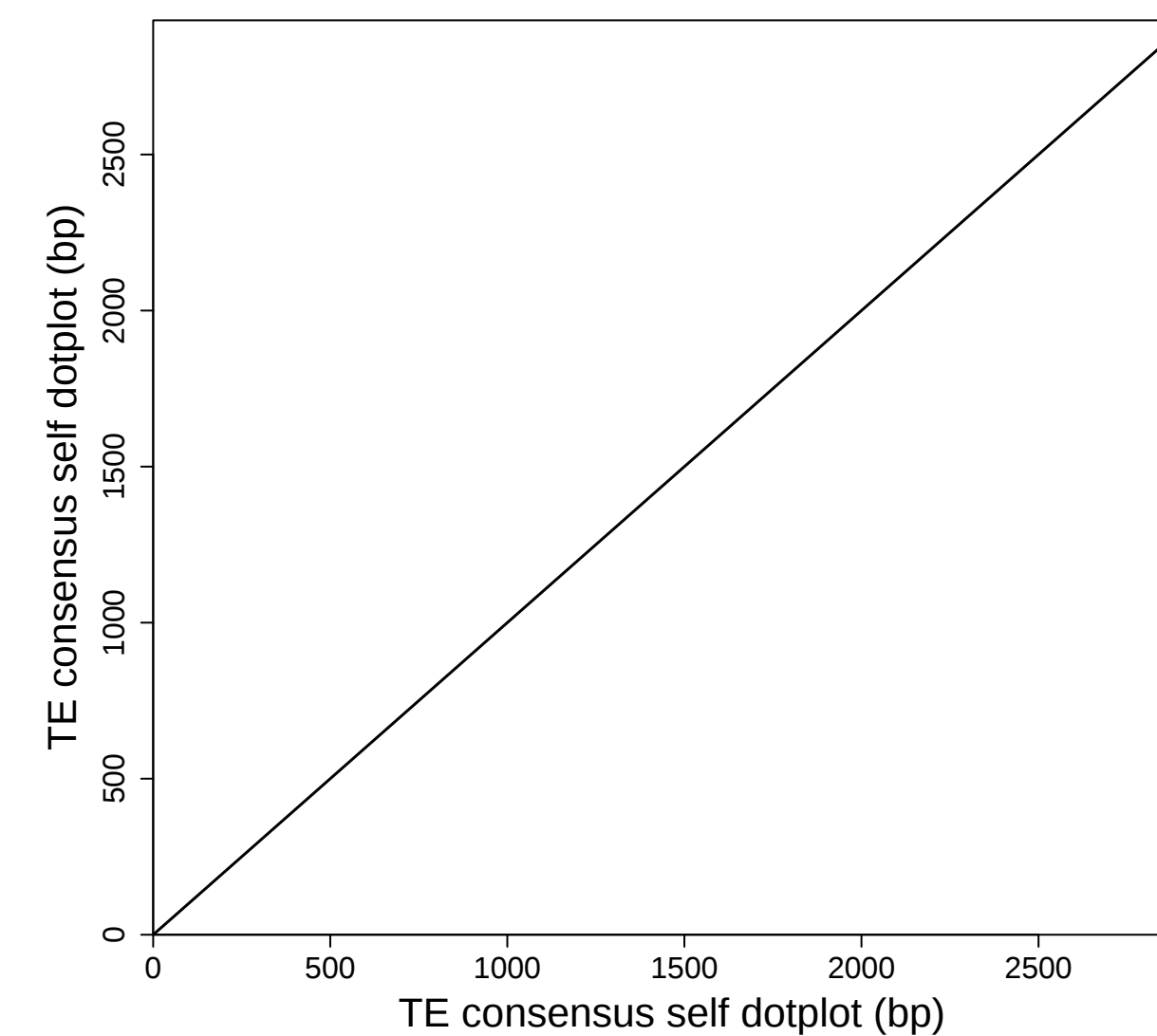
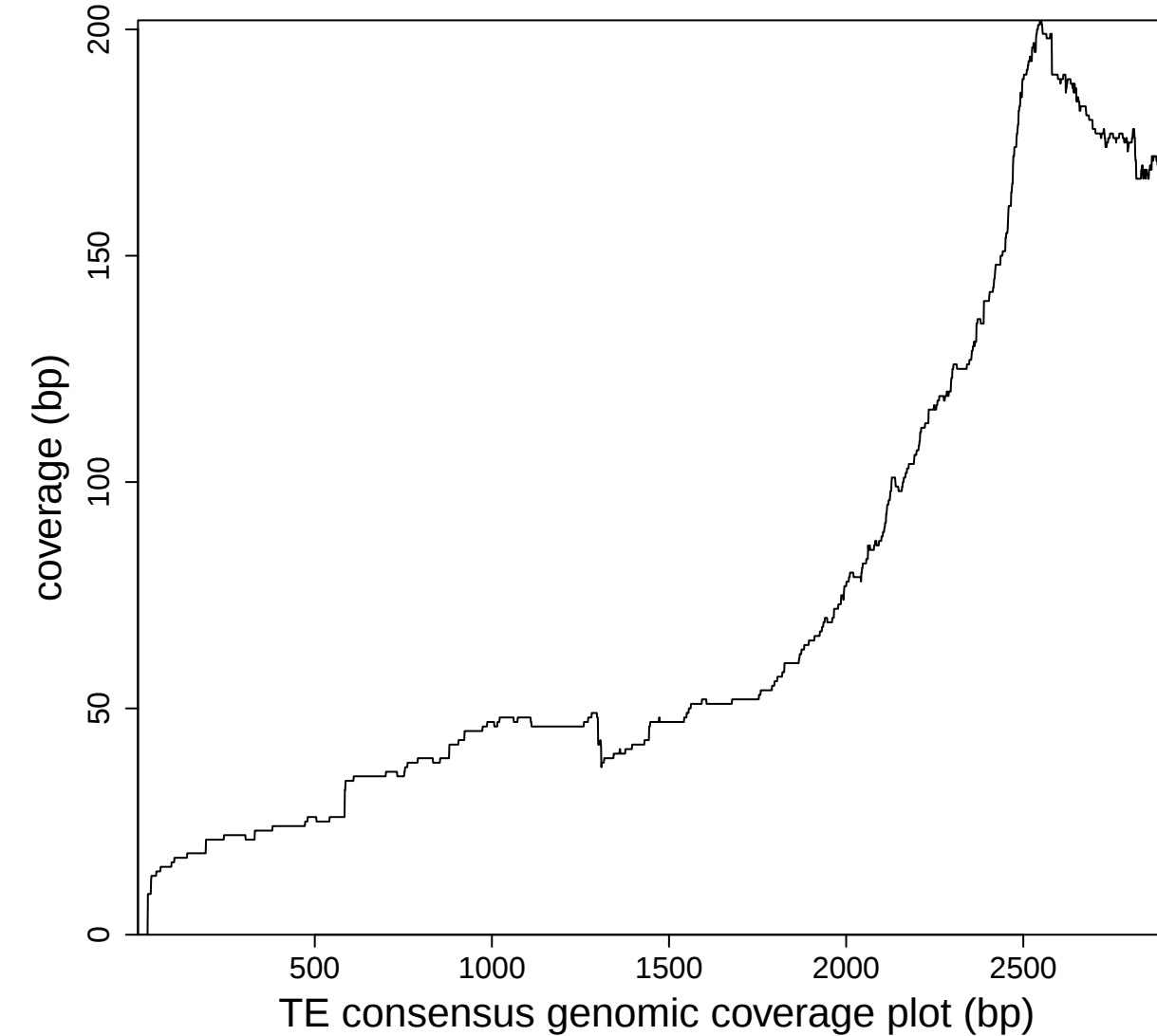
After TEtrimmer Extended plot Blue lines are boundaries



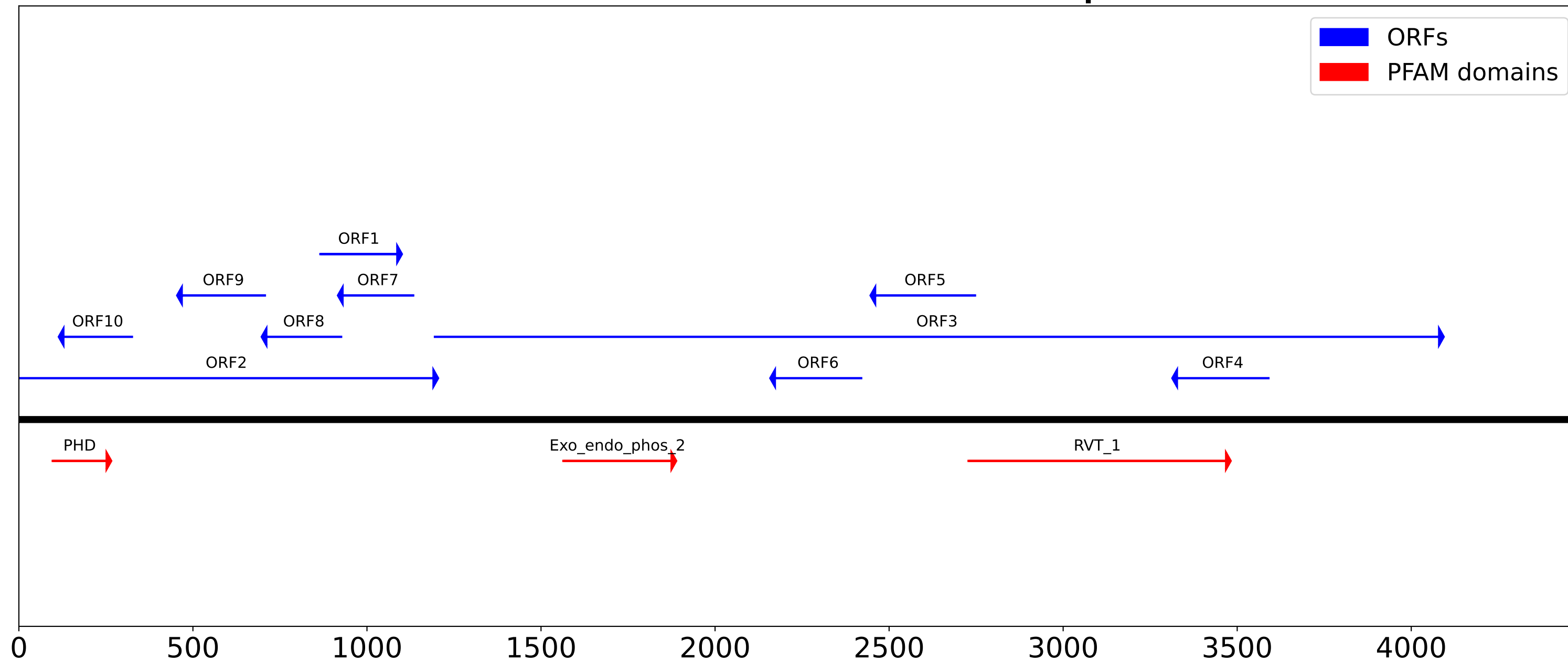
TE: rnd_5_family_383
size: 2930bp; fragments: 296; full length: 13 (>=2637bp)



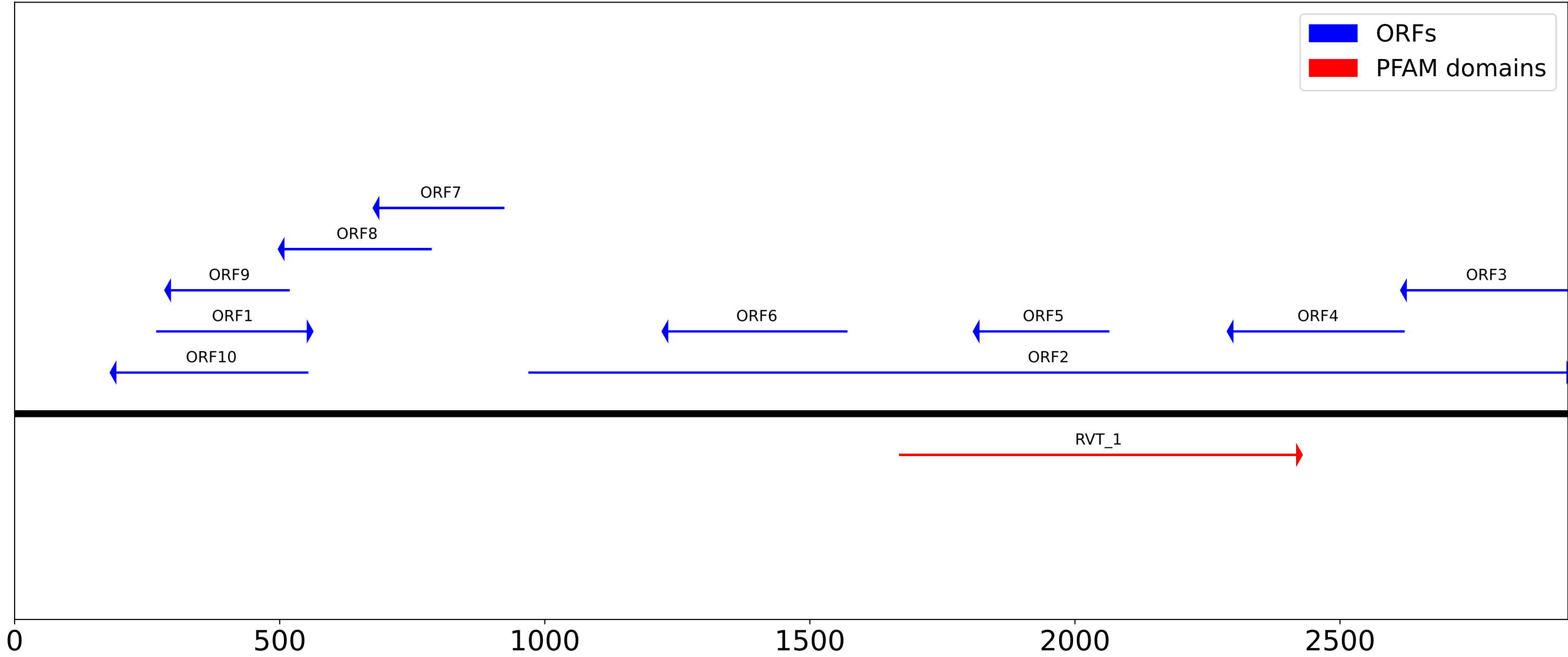
Before TEtrimmer 2930 bp



After TEtrimmer ORF and PFAM domain plot



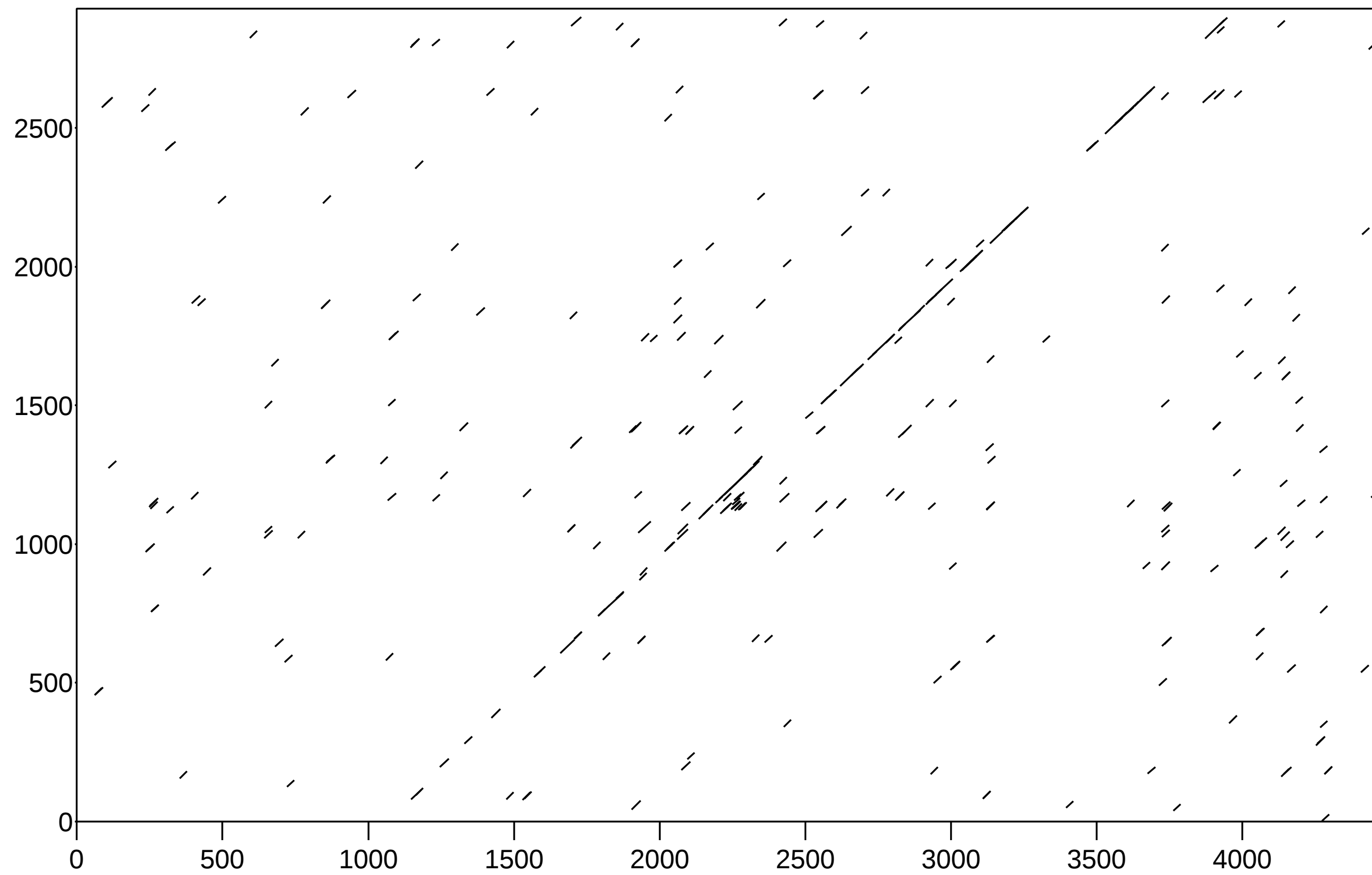
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 10/10/25)

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)